

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/050,873

DATE: 02/07/2002

TIME: 07:35:13

Input Set : A:\PZ029P2seqlist.txt

Output Set: N:\CRF3\02072002\J050873.raw

1 <110> APPLICANT: Ruben et al.

3 <120> TITLE OF INVENTION: 94 Human secreted proteins

5 <130> FILE REFERENCE: PZ029P2

7 <140> CURRENT APPLICATION NUMBER: US/10/050,873

8 <141> CURRENT FILING DATE: 2002-01-18

10 <150> PRIOR APPLICATION NUMBER: 60/263,230

11 <151> PRIOR FILING DATE: 2001-01-23

13 <150> PRIOR APPLICATION NUMBER: 60/263,681

14 <151> PRIOR FILING DATE: 2001-01-24

16 <150> PRIOR APPLICATION NUMBER: 09/461,325

17 <151> PRIOR FILING DATE: 1999-12-14

19 <150> PRIOR APPLICATION NUMBER: PCT/US99/13418

20 <151> PRIOR FILING DATE: 1999-06-15

22 <150> PRIOR APPLICATION NUMBER: 60/089,507

23 <151> PRIOR FILING DATE: 1998-06-16

25 <150> PRIOR APPLICATION NUMBER: 60/089,508

26 <151> PRIOR FILING DATE: 1998-06-16

28 <150> PRIOR APPLICATION NUMBER: 60/089,509

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31 <150> PRIOR APPLICATION NUMBER: 60/089,510

32 <151> PRIOR FILING DATE: 1998-06-16 34 <150> PRIOR APPLICATION NUMBER: 60/090,112

35 <151> PRIOR FILING DATE: 1998-06-22

37 <150> PRIOR APPLICATION NUMBER: 60/090,113

38 <151> PRIOR FILING DATE: 1998-06-22

40 <160> NUMBER OF SEQ ID NOS: 550

42 <170> SOFTWARE: PatentIn Ver. 2.0

44 <210> SEQ ID NO: 1

45 <211> LENGTH: 733

46 <212> TYPE: DNA

47 <213> ORGANISM: Homo sapiens

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60 acaagagcag gtggcagcag gggaacgtct tctcatgctc cgtgatgcat gaggctctgc

ENTERED

660

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Input Set : A:\PZ029P2seqlist.txt Output Set: N:\CRF3\02072002\J050873.raw 720 61 acaaccacta cacqcagaaq aqcctctccc tqtctccqqq taaatqaqtq cqacqqccqc 733 62 gactctagag gat 64 <210> SEQ ID NO: 2 65 <211> LENGTH: 5 66 <212> TYPE: PRT 67 <213> ORGANISM: Homo sapiens 69 <220> FEATURE: 70 <221> NAME/KEY: Site 71 <222> LOCATION: (3) 72 <223> OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids 74 <400> SEQUENCE: 2 75 Trp Ser Xaa Trp Ser 76 1 78 <210> SEQ ID NO: 3 79 <211> LENGTH: 86 80 <212> TYPE: DNA 81 <213> ORGANISM: Artificial Sequence 83 <220> FEATURE: 84 <221> NAME/KEY: Primer_Bind 85 <223> OTHER INFORMATION: Synthetic sequence with 4 tandem copies of the GAS binding site 86 found in the IRF1 promoter (Rothman et al., Immunity 1:457-468 (1994)), 18 nucleotides complementary to the SV40 early promoter, 87 and a Xho I restriction site. 90 <400> SEQUENCE: 3 91 gegeetegag attteecega aatetagatt teecegaaat gattteeceg aaatgattte 60 92 cccgaaatat ctgccatctc aattag 94 <210> SEQ ID NO: 4 95 <211> LENGTH: 27 96 <212> TYPE: DNA 97 <213> ORGANISM: Artificial Sequence 99 <220> FEATURE: 100 <221> NAME/KEY: Primer_Bind 101 <223> OTHER INFORMATION: Synthetic sequence complementary to the SV40 promter; includes a 102 Hind III restriction site. 104 <400> SEOUENCE: 4 105 gcggcaagct ttttgcaaag cctaggc 107 <210> SEQ ID NO: 5 108 <211> LENGTH: 271 109 <212> TYPE: DNA 110 <213> ORGANISM: Artificial Sequence 112 <220> FEATURE: 113 <221> NAME/KEY: Protein_Bind 114 <223> OTHER INFORMATION: Synthetic promoter for use in biological assays; includes GAS binding sites found in the IRF1 promoter (Rothman et al., Immunity 115 1:457-468 (1994)). 116 118 <400> SEQUENCE: 5 119 ctcgagattt ccccgaaatc tagatttccc cgaaatgatt tccccgaaat gatttccccg 60 120 aaatatctgc catctcaatt agtcagcaac catagtcccg cccctaactc cgcccatccc 120 121 gcccctaact ccgcccagtt ccgcccattc tccgccccat ggctgactaa ttttttttat

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Input Set : A:\PZ029P2seqlist.txt Output Set: N:\CRF3\02072002\J050873.raw 122 ttatgcagag gccgaggccg cctcggcctc tgagctattc cagaagtagt gaggaggctt 240 271 123 ttttggaggc ctaggctttt gcaaaaagct t 125 <210> SEQ ID NO: 6 126 <211> LENGTH: 32 127 <212> TYPE: DNA 128 <213> ORGANISM: Artificial Sequence 130 <220> FEATURE: 131 <221> NAME/KEY: Primer_Bind 132 <223> OTHER INFORMATION: Synthetic primer complementary to human genomic EGR-1 promoter sequence (Sakamoto et al., Oncogene 6:867-871 (1991)); includes a 133 Xho I restriction site. 134 136 <400> SEQUENCE: 6 32 137 gcgctcgagg gatgacagcg atagaacccc gg 139 <210> SEQ ID NO: 7 140 <211> LENGTH: 31 141 <212> TYPE: DNA 142 <213> ORGANISM: Artificial Sequence 144 <220> FEATURE: 145 <221> NAME/KEY: Primer_Bind 146 <223> OTHER INFORMATION: Synthetic primer complementary to human genomic EGR-1 promoter sequence (Sakamoto et al., Oncogene 6:867-871 (1991)); includes a 147 148 Hind III restriction site. 150 <400> SEQUENCE: 7 31 151 gcgaagette gcgaeteece ggateegeet e 153 <210> SEQ ID NO: 8 154 <211> LENGTH: 12 155 <212> TYPE: DNA 156 <213> ORGANISM: Homo sapiens 158 <400> SEQUENCE: 8 12 159 ggggactttc cc 161 <210> SEQ ID NO: 9 162 <211> LENGTH: 73 163 <212> TYPE: DNA 164 <213> ORGANISM: Artificial Sequence 166 <220> FEATURE: 167 <221> NAME/KEY: Primer_Bind 168 <223> OTHER INFORMATION: Synthetic primer with 4 tandem copies of the NF-KB binding site 169 (GGGGACTTTCCC), 18 nucleotides complementary to the 5' end of the 170 SV40 early promoter sequence, and a XhoI restriction site. 172 <400> SEQUENCE: 9 173 geggeetega ggggaettte eeggggaett teeggggaet tteegggaet tteeateetg 60 73 174 ccatctcaat tag 176 <210> SEQ ID NO: 10 177 <211> LENGTH: 256 178 <212> TYPE: DNA 179 <213> ORGANISM: Artificial Sequence 181 <220> FEATURE: 182 <221> NAME/KEY: Protein_Bind

183 <223> OTHER INFORMATION: Synthetic promoter for use in biological assays; includes

RAW SEQUENCE LISTING

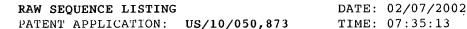
PATENT APPLICATION: US/10/050,873

NF-KB

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			cctgcagtgg				240		
			ttcacctaca				300		
			aaataccaca				360		
			aaaaagctta				420		
			cattctttgt				480		
			agctgtaaag				540		
			aaaagaagac				600		
			ctaatatgaa				660		
			gttttattag				720		
			atcctactca				780		
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			aacatcagga				240		
			ggggtcgtcg				300		
223	atazaataat	agetattata	gatgaagcgc	gacatatact	gaccaygta	ctccaacata	360		
			cgtggccgac				420		
							480		
			gaagggcgcg				540		
			ctcgggcttc						
			ggtcctgtgg				600 660		
			cttcccgggg						
			ctcatcctga				720 780		
			ccggccgggc						
			ctgccgcacc				840		
			gcagccctgc				900		
			cgaggcaagc				960		
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249 tcaacaggat attettcaag gaaaatgaac cccacactag gcctggccat ttttctggct
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250 gttctcctca cggtgaaagg tcttctaaag ccgagcttct caccaaggaa ttataaagct
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251 ttgagcgagg tccaaggatg gaagcaaagg atggcagcca aggagcttgc aaggcagaac
                                                                           300
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252 atggacttag gctttaagct gctcaagaag ctggcctttt acaaccctgg caggaacatc
253 ttcctatccc ccttgagcat ctctacagct ttctccatgc tgtgcctggg tgcccaggac
                                                                           420
254 agcaccctgg acgagatcaa gcaggggttc aacttcagaa agatgccaga aaaagatctt
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                                                                           540
256 agcattggga acacgctgtt cattgaccag aggctgcagc cacagcgtaa gtttttggaa
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257 gatqccaaqa acttttacag tgccgaaacc atccttacca actttcagaa tttggaaatg
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264 tocagatgga aaacattact gtcacgcagg gtcgtagacg tgtctgtacc cagactccac
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265 atgacgggca cettegacet gaagaagact eteteetaca taggtgtete caaaatettt
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266 gaggaacatg gtgatctcac caagatcgcc cctcatcgca gcctgaaagt gggcgaggct
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267 gtgcacaagg ctgagctgaa gatggatgag aggggtacgg aaggggccgc tggcaccgga
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268 gcacagacte tgeccatgga gacaceaete gtegteaaga tagacaaace etatetgetg
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269 ctgatttaca gcgagaaaat accttccgtg ctcttcctgg gaaagattgt taaccctatt
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270 ggaaaataaa ggagaattee tgettgeeae agaeeeegaa aaaaaaaaa aaaaagggeg
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291 <223> OTHER INFORMATION: n equals a,t,g, or c
293 <220> FEATURE:
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Use of n and/or Xaa has been detected in the Scquence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

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Input Set : A:\PZ029P2seqlist.txt
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L:536 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
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L:2141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64
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